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Applicant(s): Dominic E. Cosgrove

Group Art Unit:

Serial No.:

09/970,318

Examirer:

TECH UENTER 1600/2900 Bao Thuy L. Nguyen

Confirmation No.: 1885

October 3, 2001

For:

Filed:

IMMUNODIAGNOSTIC DETERMINATION OF USHER SYNDROME

TYPE HA

## PRELIMINARY AMENDMENT

**Assistant Commissioner for Patents** P.O. Box 2327 Arlington, VA 22202

Dear Sir:

Prior to taking up the above-identified application for examination, please amend the application as follows:

## In the Specification

Please replace the paragraph beginning at page 2, line 1, with the following rewritten paragraph. Per 37 C.F.R. §1.121, this paragraph is also shown in Appendix A with notations to indicate the changes made.

Usher syndrome Type II is the most common of the three Usher syndromes. Although originally it was believed that Usher Type II accounted for only about 10% of all Usher cases, more recent research shows that Type II actually accounts for over half of all Usher cases. The USH2A gene has been localized to chromosome 1q41 between D1S474 and AFM144FX2 (Kimberling et al., Am. J. Hum. Genet., 56:216-223 (1995); Sumegi et al., Genomics, 35:79-86 (1996)), and more recently, the gene has been identified (Eudy et al., Science, 280:1753-1757 (1998)). However, there are Usher Type II families whose disease locus cannot be linked to the 1q41 region. Two new Usher II loci have been localized to 3p and 5q (Pieke-Dahl et al., J. Med. Genet., 37:256-262 (2000);

